

# REPLACEMENT SHEET

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Title: PEPTIDE SEQUENCES SPECIFIC FOR THE HEPATIC STAGES OF P. FALCIPARUM BEARING EPITOPES CAPABLE OF STIMULATING THE T LYMPHOCYTES

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Sheet 1 of 4

LSN.3'STOP -> 1-phase Translation

DNA sequence 1482 b.p. CAAGAACAACAA ... ATGAAACTATAA linear

1 / 1	31 / 11	
CAA GAA CAA CAA AGC GAT CTA GAA CAA GAG AGA CGT GCT AAA GAA AAG TTG CAA GAA GAA		
gln glu gln gln ser asp leu glu gln gln glu arg ala lys glu lys leu gln glu gln		
61 / 21	91 / 31	
CAA AGC GAT TTA GAA CAA CAA CTT GCT AAA GAA AAG TTA CAA GAG CAG CAA AGC GAT		
gln ser asp leu glu gln asp arg leu ala lys glu lys leu gln glu gln gln ser asp		
121 / 41	151 / 51	
TTA GAA CAA GAG AGA CTT GCT AAA GAA AAG TTG CAA GAA CAA CAA AGC GAT CTA GAA CAA		
leu glu gln glu arg leu ala lys glu lys leu gln glu gln gln ser asp leu glu gln		
181 / 61	211 / 71	
GAG AGA CGT GCT AAA GAA AAG TTG CAA GAA CAA CAA AGC GAT TTA GAA CAA GAG AGA CGT		
glu arg arg ala lys glu lys leu gln glu gln gln ser asp leu glu gln glu arg		
241 / 81	271 / 91	
GCT AAA GAA AAG TTG CAA GAA CAA CAA AGC GAT TTA GAA CAA GAT AGA CTT GCT AAA GAA		
ala lys glu lys leu gln glu gln gln ser asp leu glu gln asp arg leu ala lys glu		
301 / 101	331 / 111	
AAG TTA CAA GAG CAG CAA AGC GAT TTA GAA CAA GAG AGA CGT GCT AAA GAA AAG TTG CAA		
lys leu gln glu gln ser asp leu glu gln glu arg arg ala lys glu lys leu gln		

FIGURE 10A

361 /	GAA CAA	AGC GAT	TTA GAA	AGA CGT	GCT GCT	AAA GAA	AAG AAG	TTG CAA	GAA GAA	CAA CAA	GAT CAA	CAA CAA	gln gln
421 /	gln gln	ser asp	leu leu	arg arg	ala ala	lys lys	lys lys	leu leu	gln gln	gln gln	asp asp	gln gln	gln
481 /	ser asp	leu leu	gln gln	gln gln	leu leu	ala ala	lys lys	leu leu	gln gln	gln gln	ser ser	asp asp	leu leu
541 /	GAA CAA	GAG AGA	GCT GCT	AAA GAA	AAG TTT	GCT GCT	AAA GAA	AAG TTT	CAA CAA	GAT CAA	AGC GAT	CAA CAA	TTA TTA
601 /	arg arg	ala ala	lys lys	gln gln	lys lys	leu leu	gln gln	gln gln	ser ser	asp asp	leu leu	gln gln	gln
661 /	AAA GAA	AAG TTT	GCT GCT	AAA GAA	AAG TTT	GCT GCT	AAA GAA	AAG TTT	CAA CAA	GAT CAA	AGC GAT	CAA CAA	GAT CAA
721 /	asn leu	lys lys	leu leu	gln gln	lys lys	leu leu	gln gln	gln gln	ser ser	asp asp	leu leu	gln gln	gln
781 /	GAA ATA	CCA GCT	ATA GAA	AAG TTT	GCT GCT	AAA GAA	AAG TTT	GCT GCT	AAA GAA	AAG TTT	GCT GCT	CAA CAA	CAA CAA
841 /	gln gln	ser asp	leu leu	gln gln	lys lys	leu leu	gln gln	gln gln	ser ser	asp asp	leu leu	gln gln	gln
901 /	TCT TCT	TTA CCT	CAG GAC	AAC AAC	AGA GGG	AAT AGT	AGA GGG	AAT AGT	TCC TCC	AAG GAA	ATG ATA	TCT ATA	ATA ATA
961 /	ser ser	leu leu	pro pro	gln gln	asp asp	asn asn	arg arg	gly gly	lys lys	gln gln	met met	ser ser	ile ile

FIGURE 10B

841 / 281	GAA AAA ACA AAT AGA GAA TCT ATT ACA	871 / 291	GAA GGA CGA AGG GAT ATA CAT AAA
901 / 301	glu lys thr asn arg glu ser ile thr		glu gly arg asp ile his lys
GGA CAT CTT GAA AAG GAA AAG GAT GGT	931 / 311		
gly his leu glu lys lys asp gly ser ile		ATA AAA CCA GAA AAA GAA GAT AAA TCT	
961 / 321	991 / 331		lys glu asp lys ser
GCT GAC ATA CAA AAT CAT ACA TTA GAG		ATT TCT GAT GTT AAT GAT TTT CAA	
ala asp ile gln asn his thr leu glu thr	1051 / 351		asn val asp phe gln
1021 / 341			
ATA AGT AAG TAT GAG GAT GAA ATA AGT		GAC GAT TCA TTA ATA GAT GAA GAA	
ile ser lys tyr glu asp glu ile ser ala		glu tyr asp ser leu ile asp glu glu	
1081 / 361	1111 / 371		
GAA GAT GAT GAA GAC TTA GAC GAA TTT		AGG CCT ATT GTG CAA TAT GAC AAT TTC CAA GAT	
glu asp asp glu asp leu asp glu phe		lys pro ile val gln tyr asp phe gln asp	
1141 / 381	1171 / 391		
GAA GAA AAC ATA GGA ATT TAT AAA GAA		CTA GAA GAT TTG ATA GAG AAA AAT GAA AAT TTA	
glu glu asn ile gly ile tyr lys glu leu		glu asp leu ile glu lys asn glu asn leu	
1201 / 401	1231 / 411		
GAT GAT TTA GAT GAA GGA ATA GAA AAA		TCA TCA TCT GAA GAA AAA ATA AAA	
asp asp leu asp glu gly ile glu lys ser		glu leu ser glu glu lys ile lys	
1261 / 421	1291 / 431		
AAA GGA AAG AAA TAT GAA AAA ACA AAG		GAT AAT TTT AAA CCA AAT GAT AAA AGT TTG	
lys gly lys lys tyr glu lys thr lys		asn phe lys pro asn asp lys ser leu	

FIGURE 10C

1321 / 441	1351 / 451
TAT GAT GAG CAT ATT AAA AAT GAT AAG CAG GTT AAT AAG GAA AAG GAA AAA	
tyr asp glu his ile lys lys tyr lys asn asp lys gln val asn lys glu lys glu lys	
1381 / 461	1411 / 471
TTC ATA AAA TCA TTG TTT CAT ATA TTT GAC GGA GAC AAT GAA ATT TTA CAG ATC GTG GAT	
phe ile lys ser leu phe his ile phe asp gly asp asn glu ile leu gln ile val asp	
1441 / 481	1471 / 491
GAG TTA TCT GAA GAT ATA ACT AAA TAT TTT ATG AAA CTA TAA AAG GTT ATA TAT	
glu leu ser glu asp ile thr lys tyr phe met lys leu OCH lys val ile tyr	

FIGURE 10D